

SCORE Search Results Details for Application 10552515 and Search Result 20080630_144103_us-10-552-515-10.rai.

[Score Home Page](#)

[Retrieve Application List](#)

[SCORE System Overview](#)

[SCORE FAQ](#)

[Comments / Suggestions](#)

This page gives you Search Results detail for the Application 10552515 and Search Result 20080630_144103_us-10-552-515-10.rai.

[Go Back to previous page](#)

GenCore version 6.2.1
Copyright (c) 1993 - 2008 Biocceleration Ltd.

OM protein - protein search, using sw model

Run on: June 30, 2008, 17:46:21 ; Search time 40 Seconds
(without alignments)
42.303 Million cell updates/sec

Title: US-10-552-515-10
Perfect score: 44
Sequence: 1 KIYVSLAHV 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1143754 seqs, 186252778 residues

Total number of hits satisfying chosen parameters: 1143754

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:
1: /ABSS/Data/CRF/ptodata/1/iaa/5_COMB.pep:
2: /ABSS/Data/CRF/ptodata/1/iaa/6_COMB.pep:
3: /ABSS/Data/CRF/ptodata/1/iaa/7_COMB.pep:
4: /ABSS/Data/CRF/ptodata/1/iaa/H_COMB.pep:
5: /ABSS/Data/CRF/ptodata/1/iaa/PCTUS_COMB.pep:
6: /ABSS/Data/CRF/ptodata/1/iaa/RE_COMB.pep:
7: /ABSS/Data/CRF/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	34	77.3	389	3	US-10-369-493-10502	Sequence 10502, A
2	33	75.0	110	3	US-10-703-032-129843	Sequence 129843,
3	33	75.0	566	1	US-08-666-367B-5	Sequence 5, Appli
4	33	75.0	566	2	US-09-143-438-5	Sequence 5, Appli
5	32	72.7	292	3	US-10-703-032-107936	Sequence 107936,
6	32	72.7	527	3	US-10-369-493-7810	Sequence 7810, Ap
7	32	72.7	1414	3	US-10-667-891-2	Sequence 2, Appli
8	31	70.5	77	3	US-10-703-032-144572	Sequence 144572,
9	31	70.5	103	2	US-09-248-796A-26386	Sequence 26386, A
10	31	70.5	212	3	US-10-703-032-174397	Sequence 174397,
11	31	70.5	284	2	US-09-252-991A-29913	Sequence 29913, A
12	31	70.5	483	2	US-09-106-194-4	Sequence 4, Appli
13	31	70.5	519	2	US-09-949-016-9399	Sequence 9399, Ap
14	31	70.5	525	2	US-09-773-426A-1	Sequence 1, Appli
15	31	70.5	525	2	US-10-314-881-1	Sequence 1, Appli
16	31	70.5	525	2	US-09-495-823-1	Sequence 1, Appli
17	31	70.5	525	3	US-10-426-776-10	Sequence 10, Appli
18	31	70.5	525	3	US-10-123-292-56	Sequence 56, Appli
19	31	70.5	525	3	US-10-152-398-56	Sequence 56, Appli
20	31	70.5	525	3	US-10-123-907-56	Sequence 56, Appli
21	31	70.5	525	3	US-10-147-512-56	Sequence 56, Appli
22	31	70.5	525	3	US-10-147-485-56	Sequence 56, Appli
23	31	70.5	525	3	US-10-124-814-56	Sequence 56, Appli
24	31	70.5	525	3	US-10-124-822-56	Sequence 56, Appli
25	31	70.5	525	3	US-10-131-833A-56	Sequence 56, Appli
26	31	70.5	525	3	US-10-142-419-56	Sequence 56, Appli
27	31	70.5	525	3	US-10-152-375-56	Sequence 56, Appli
28	31	70.5	525	3	US-10-131-818A-56	Sequence 56, Appli
29	31	70.5	525	3	US-10-145-873-56	Sequence 56, Appli
30	31	70.5	525	3	US-10-152-395-56	Sequence 56, Appli
31	31	70.5	525	3	US-10-131-822A-56	Sequence 56, Appli
32	31	70.5	525	3	US-10-142-763-56	Sequence 56, Appli
33	31	70.5	525	3	US-10-128-694A-56	Sequence 56, Appli
34	31	70.5	525	3	US-10-123-213-56	Sequence 56, Appli
35	31	70.5	525	3	US-10-123-909-56	Sequence 56, Appli
36	31	70.5	525	3	US-10-131-826A-56	Sequence 56, Appli
37	31	70.5	525	3	US-10-147-513-56	Sequence 56, Appli
38	31	70.5	525	3	US-10-121-043-56	Sequence 56, Appli
39	31	70.5	525	3	US-10-139-980-56	Sequence 56, Appli
40	31	70.5	525	3	US-10-131-819A-56	Sequence 56, Appli
41	31	70.5	525	3	US-10-123-212-56	Sequence 56, Appli
42	31	70.5	525	3	US-10-131-813A-56	Sequence 56, Appli
43	31	70.5	525	3	US-10-140-021-56	Sequence 56, Appli
44	31	70.5	525	3	US-10-137-869A-56	Sequence 56, Appli
45	31	70.5	525	3	US-10-140-923-56	Sequence 56, Appli

ALIGNMENTS

RESULT 1

US-10-369-493-10502

; Sequence 10502, Application US/10369493

; Patent No. 7314974
 ; GENERAL INFORMATION:
 ; APPLICANT: Cao, Yongwei
 ; APPLICANT: Hinkle, Gregory J.
 ; APPLICANT: Slater, Steven C.
 ; APPLICANT: Goldman, Barry S.
 ; APPLICANT: Chen, Xianfeng
 ; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
 ; TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
 ; FILE REFERENCE: 38-10(52052)B
 ; CURRENT APPLICATION NUMBER: US/10/369,493
 ; CURRENT FILING DATE: 2003-02-28
 ; PRIOR APPLICATION NUMBER: US 60/360,039
 ; PRIOR FILING DATE: 2002-02-21
 ; NUMBER OF SEQ ID NOS: 47374
 ; SEQ ID NO 10502
 ; LENGTH: 389
 ; TYPE: PRT
 ; ORGANISM: Sphingomonas aromaticivorans
 US-10-369-493-10502

Query Match 77.3%; Score 34; DB 3; Length 389;
 Best Local Similarity 66.7%; Pred. No. 81;
 Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy	1	KIYVSLAHV 9
		: :
Db	159	KIWTSLAHI 167

RESULT 2
 US-10-703-032-129843
 ; Sequence 129843, Application US/10703032
 ; Patent No. 7214786
 ; GENERAL INFORMATION:
 ; APPLICANT: Kovalic, David K.
 ; APPLICANT: Andersen, Scott E.
 ; APPLICANT: Byrum, Joseph R.
 ; APPLICANT: Conner, Timothy W.
 ; APPLICANT: Cao, Yongwei
 ; APPLICANT: Masucci, James D.
 ; APPLICANT: Zhou, Yihua
 ; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
 ; TITLE OF INVENTION: Plants
 ; FILE REFERENCE: 38-21(53374)B
 ; CURRENT APPLICATION NUMBER: US/10/703,032
 ; CURRENT FILING DATE: 2003-11-06
 ; PRIOR APPLICATION NUMBER: 10/020,338
 ; PRIOR FILING DATE: 2001-12-12
 ; NUMBER OF SEQ ID NOS: 211164
 ; SEQ ID NO 129843
 ; LENGTH: 110
 ; TYPE: PRT
 ; ORGANISM: Triticum aestivum
 ; FEATURE:

;
 NAME/KEY: unsure
 LOCATION: (1)..(110)
 OTHER INFORMATION: unsure at all Xaa locations
 FEATURE:
 OTHER INFORMATION: Clone ID: PAT_TA_24261.pep
 US-10-703-032-129843

Query Match 75.0%; Score 33; DB 3; Length 110;
 Best Local Similarity 66.7%; Pred. No. 33;
 Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9
 | :| |||
 Db 6 KXYISAAHV 14

RESULT 3

US-08-666-367B-5

;
 Sequence 5, Application US/08666367B
 Patent No. 5854042
 GENERAL INFORMATION:
 APPLICANT: Shuichi TSUJI et al.
 TITLE OF INVENTION: NOVEL SUGAR-CHAIN SYNTHETASE AND PROCESS FOR
 TITLE OF INVENTION: PRODUCING THE SAME
 NUMBER OF SEQUENCES: 8
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Wenderoth, Lind & Ponack
 STREET: 805 Fifteenth Street, N.W., #700
 CITY: Washington
 STATE: D.C.
 COUNTRY: U.S.A.
 ZIP: 20005
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb
 COMPUTER: IBM Compatible
 OPERATING SYSTEM: MS-DOS
 SOFTWARE: Wordperfect 5.1
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/666,367B
 FILING DATE: August 19, 1996
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER:
 FILING DATE:
 ATTORNEY/AGENT INFORMATION:
 NAME: Warren M. Cheek, Jr.
 REGISTRATION NUMBER: 33,367
 REFERENCE/DOCKET NUMBER:
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 202-371-8850
 TELEFAX:
 TELEX:
 INFORMATION FOR SEQ ID NO: 5:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 566 amino acids
 TYPE: amino acid

;
; STRANDEDNESS: single

;
; TOPOLOGY: linear

US-08-666-367B-5

Query Match 75.0%; Score 33; DB 1; Length 566;
Best Local Similarity 66.7%; Pred. No. 2e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9

||| |||:

Db 46 KIYQSIAHM 54

RESULT 4

US-09-143-438-5

;
; Sequence 5, Application US/09143438

;
; Patent No. 6218161

;
; GENERAL INFORMATION:

;
; APPLICANT: Shuichi TSUJI et al.

;
; TITLE OF INVENTION: NOVEL SUGAR-CHAIN SYNTHETASE AND PROCESS FOR

;
; TITLE OF INVENTION: PRODUCING THE SAME

;
; NUMBER OF SEQUENCES: 8

;
; CORRESPONDENCE ADDRESS:

;
; ADDRESSEE: Wenderoth, Lind & Ponack, L.L.P.

;
; STREET: 2033 K Street, N.W., #800

;
; CITY: Washington

;
; STATE: D.C.

;
; COUNTRY: U.S.A.

;
; ZIP: 20006

;
; COMPUTER READABLE FORM:

;
; MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb

;
; COMPUTER: IBM Compatible

;
; OPERATING SYSTEM: MS-DOS

;
; SOFTWARE: Wordperfect 5.1

;
; CURRENT APPLICATION DATA:

;
; APPLICATION NUMBER: US/09/143,438

;
; FILING DATE: August 28, 1998

;
; CLASSIFICATION:

;
; PRIOR APPLICATION DATA:

;
; APPLICATION NUMBER: 08/666,367

;
; FILING DATE: August 19, 1996

;
; ATTORNEY/AGENT INFORMATION:

;
; NAME: Warren M. Cheek, Jr.

;
; REGISTRATION NUMBER: 33,367

;
; REFERENCE/DOCKET NUMBER:

;
; TELECOMMUNICATION INFORMATION:

;
; TELEPHONE: 202-721-8200

;
; TELEFAX: 202-721-8250

;
; TELEX:

;
; INFORMATION FOR SEQ ID NO: 5:

;
; SEQUENCE CHARACTERISTICS:

;
; LENGTH: 566 amino acids

;
; TYPE: amino acid

;
; STRANDEDNESS: single

;
; TOPOLOGY: linear

US-09-143-438-5

Query Match 75.0%; Score 33; DB 2; Length 566;
 Best Local Similarity 66.7%; Pred. No. 2e+02;
 Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9
 !!! |:||:
 Db 46 KIYQSIAHM 54

RESULT 5

US-10-703-032-107936

; Sequence 107936, Application US/10703032
 ; Patent No. 7214786
 ; GENERAL INFORMATION:
 ; APPLICANT: Kovalic, David K.
 ; APPLICANT: Andersen, Scott E.
 ; APPLICANT: Byrum, Joseph R.
 ; APPLICANT: Conner, Timothy W.
 ; APPLICANT: Cao, Yongwei
 ; APPLICANT: Masucci, James D.
 ; APPLICANT: Zhou, Yihua
 ; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
 ; TITLE OF INVENTION: Plants
 ; FILE REFERENCE: 38-21(53374)B
 ; CURRENT APPLICATION NUMBER: US/10/703,032
 ; CURRENT FILING DATE: 2003-11-06
 ; PRIOR APPLICATION NUMBER: 10/020,338
 ; PRIOR FILING DATE: 2001-12-12
 ; NUMBER OF SEQ ID NOS: 211164
 ; SEQ ID NO 107936
 ; LENGTH: 292
 ; TYPE: PRT
 ; ORGANISM: Triticum aestivum
 ; FEATURE:
 ; OTHER INFORMATION: Clone ID: PAT_TA_2354.pep
 US-10-703-032-107936

Query Match 72.7%; Score 32; DB 3; Length 292;
 Best Local Similarity 75.0%; Pred. No. 1.6e+02;
 Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KIYVSLAH 8
 ! ||:|||:
 Db 246 KAYVTLAH 253

RESULT 6

US-10-369-493-7810

; Sequence 7810, Application US/10369493
 ; Patent No. 7314974
 ; GENERAL INFORMATION:
 ; APPLICANT: Cao, Yongwei
 ; APPLICANT: Hinkle, Gregory J.
 ; APPLICANT: Slater, Steven C.

; APPLICANT: Goldman, Barry S.
 ; APPLICANT: Chen, Xianfeng
 ; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
 ; TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
 ; FILE REFERENCE: 38-10(52052)B
 ; CURRENT APPLICATION NUMBER: US/10/369,493
 ; CURRENT FILING DATE: 2003-02-28
 ; PRIOR APPLICATION NUMBER: US 60/360,039
 ; PRIOR FILING DATE: 2002-02-21
 ; NUMBER OF SEQ ID NOS: 47374
 ; SEQ ID NO 7810
 ; LENGTH: 527
 ; TYPE: PRT
 ; ORGANISM: Rhodobacter sphaeroides
 US-10-369-493-7810

Query Match 72.7%; Score 32; DB 3; Length 527;
 Best Local Similarity 50.0%; Pred. No. 3e+02;
 Matches 4; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IYVSLAHV 9
 ::|:||:|:
 Db 304 VYISMALH 311

RESULT 7

US-10-667-891-2

; Sequence 2, Application US/10667891
 ; Patent No. 7067259
 ; GENERAL INFORMATION:
 ; APPLICANT: ROTH, CHARLES W.
 ; APPLICANT: BREY, PAUL T.
 ; APPLICANT: HOLM, INGE
 ; APPLICANT: GRAILLES, MARINE
 ; APPLICANT: RZHETSKY, ANDREY
 ; TITLE OF INVENTION: MULTIDRUG RESISTANCE PROTEINS IN DROSOPHILA AND
 ; TITLE OF INVENTION: ANOPHELES
 ; FILE REFERENCE: 03495.0294-00000
 ; CURRENT APPLICATION NUMBER: US/10/667,891
 ; CURRENT FILING DATE: 2003-09-23
 ; PRIOR APPLICATION NUMBER: 60/413,469
 ; PRIOR FILING DATE: 2002-09-26
 ; NUMBER OF SEQ ID NOS: 76
 ; SOFTWARE: PatentIn Ver. 3.2
 ; SEQ ID NO 2
 ; LENGTH: 1414
 ; TYPE: PRT
 ; ORGANISM: Anopheles gambiae
 US-10-667-891-2

Query Match 72.7%; Score 32; DB 3; Length 1414;
 Best Local Similarity 71.4%; Pred. No. 8.7e+02;
 Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 3 YVSLAHV 9

|:|:|:|

Db 1248 YISIAHV 1254

RESULT 8

US-10-703-032-144572

; Sequence 144572, Application US/10703032

; Patent No. 7214786

; GENERAL INFORMATION:

; APPLICANT: Kovalic, David K.

; APPLICANT: Andersen, Scott E.

; APPLICANT: Byrum, Joseph R.

; APPLICANT: Conner, Timothy W.

; APPLICANT: Cao, Yongwei

; APPLICANT: Masucci, James D.

; APPLICANT: Zhou, Yihua

; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With

; TITLE OF INVENTION: Plants

; FILE REFERENCE: 38-21(53374)B

; CURRENT APPLICATION NUMBER: US/10/703,032

; CURRENT FILING DATE: 2003-11-06

; PRIOR APPLICATION NUMBER: 10/020,338

; PRIOR FILING DATE: 2001-12-12

; NUMBER OF SEQ ID NOS: 211164

; SEQ ID NO 144572

; LENGTH: 77

; TYPE: PRT

; ORGANISM: Triticum aestivum

; FEATURE:

; OTHER INFORMATION: Clone ID: PAT_TA_38990.pep

US-10-703-032-144572

Query Match 70.5%; Score 31; DB 3; Length 77;

Best Local Similarity 75.0%; Pred. No. 59;

Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 2 IYVSLAHV 9

|||| ||:

Db 15 IYVSHAI 22

RESULT 9

US-09-248-796A-26386

; Sequence 26386, Application US/09248796A

; Patent No. 6747137

; GENERAL INFORMATION:

; APPLICANT: Keith Weinstock et al

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS

; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: 107196.132

; CURRENT APPLICATION NUMBER: US/09/248,796A

; CURRENT FILING DATE: 1999-02-12

; PRIOR APPLICATION NUMBER: US 60/074,725

; PRIOR FILING DATE: 1998-02-13

; PRIOR APPLICATION NUMBER: US 60/096,409

; PRIOR FILING DATE: 1998-08-13

; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 26386
; LENGTH: 103
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-26386

Query Match 70.5%; Score 31; DB 2; Length 103;
Best Local Similarity 55.6%; Pred. No. 81;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9
Db 48 KIYISSIHI 56

RESULT 10
US-10-703-032-174397
; Sequence 174397, Application US/10703032
; Patent No. 7214786
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Andersen, Scott E.
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Conner, Timothy W.
; APPLICANT: Cao, Yongwei
; APPLICANT: Masucci, James D.
; APPLICANT: Zhou, Yihua
; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR APPLICATION NUMBER: 10/020,338
; PRIOR FILING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 174397
; LENGTH: 212
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_TA_68815.pep
US-10-703-032-174397

Query Match 70.5%; Score 31; DB 3; Length 212;
Best Local Similarity 66.7%; Pred. No. 1.8e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9
Db 69 KILVSLGHI 77

RESULT 11
US-09-252-991A-29913

; Sequence 29913, Application US/09252991A
 ; Patent No. 6551795
 ; GENERAL INFORMATION:
 ; APPLICANT: Marc J. Rubenfield et al.
 ; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
 ; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
 ; FILE REFERENCE: 107196.136
 ; CURRENT APPLICATION NUMBER: US/09/252,991A
 ; CURRENT FILING DATE: 1999-02-18
 ; PRIOR APPLICATION NUMBER: US 60/074,788
 ; PRIOR FILING DATE: 1998-02-18
 ; PRIOR APPLICATION NUMBER: US 60/094,190
 ; PRIOR FILING DATE: 1998-07-27
 ; NUMBER OF SEQ ID NOS: 33142
 ; SEQ ID NO 29913
 ; LENGTH: 284
 ; TYPE: PRT
 ; ORGANISM: Pseudomonas aeruginosa
 US-09-252-991A-29913

Query Match 70.5%; Score 31; DB 2; Length 284;
 Best Local Similarity 66.7%; Pred. No. 2.5e+02;
 Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy	1	KIYVSLAHV	9
		: :	
Db	236	RIYVNEAHV	244

RESULT 12

US-09-106-194-4

; Sequence 4, Application US/09106194
 ; Patent No. 6262234
 ; GENERAL INFORMATION:
 ; APPLICANT: Holloway, James
 ; APPLICANT: Jelinek, Laura
 ; APPLICANT: Durnam, Diane
 ; APPLICANT: Blumberg, Hal
 ; TITLE OF INVENTION: NOVEL NUCLEAR RECEPTOR POLYPEPTIDE
 ; TITLE OF INVENTION: ZPPAR4
 ; NUMBER OF SEQUENCES: 11
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: ZymoGenetics, Inc.
 ; STREET: 1201 Eastlake Avenue East
 ; CITY: Seattle
 ; STATE: WA
 ; COUNTRY: USA
 ; ZIP: 98102
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Diskette
 ; COMPUTER: IBM Compatible
 ; OPERATING SYSTEM: DOS
 ; SOFTWARE: FastSEQ for Windows Version 2.0
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/09/106,194
 ; FILING DATE:

;
 CLASSIFICATION:
 ; PRIOR APPLICATION DATA:
 ; APPLICATION NUMBER:
 ; FILING DATE:
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: Leith, Debra K
 ; REGISTRATION NUMBER: 32,619
 ; REFERENCE/DOCKET NUMBER: 96-11
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: 206-442-6674
 ; TELEFAX: 206-442-6678
 ; TELEX:
 ; INFORMATION FOR SEQ ID NO: 4:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 483 amino acids
 ; TYPE: amino acid
 ; STRANDEDNESS: single
 ; TOPOLOGY: linear
 ; MOLECULE TYPE: peptide
 US-09-106-194-4

Query Match 70.5%; Score 31; DB 2; Length 483;
 Best Local Similarity 66.7%; Pred. No. 4.4e+02;
 Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9
 ||| :| ||
 Db 411 KIYFALQHV 419

RESULT 13
 US-09-949-016-9399
 ; Sequence 9399, Application US/09949016
 ; Patent No. 6812339
 ; GENERAL INFORMATION:
 ; APPLICANT: VENTER, J. Craig et al.
 ; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
 ; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
 ; FILE REFERENCE: CL001307
 ; CURRENT APPLICATION NUMBER: US/09/949,016
 ; CURRENT FILING DATE: 2000-04-14
 ; PRIOR APPLICATION NUMBER: 60/241,755
 ; PRIOR FILING DATE: 2000-10-20
 ; PRIOR APPLICATION NUMBER: 60/237,768
 ; PRIOR FILING DATE: 2000-10-03
 ; PRIOR APPLICATION NUMBER: 60/231,498
 ; PRIOR FILING DATE: 2000-09-08
 ; NUMBER OF SEQ ID NOS: 207012
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO 9399
 ; LENGTH: 519
 ; TYPE: PRT
 ; ORGANISM: Human
 US-09-949-016-9399

Query Match 70.5%; Score 31; DB 2; Length 519;
 Best Local Similarity 66.7%; Pred. No. 4.7e+02;
 Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KIYVSLAHV 9
 ||| :| ||
 Db 447 KIYFALQHV 455

RESULT 14

US-09-773-426A-1

; Sequence 1, Application US/09773426A
 ; Patent No. 6534302
 ; GENERAL INFORMATION:
 ; APPLICANT: Glucksman, Maria Alexandra
 ; APPLICANT: Williamson, Mark
 ; APPLICANT: Tsia, Fong-Ying
 ; APPLICANT: Rudolph-Owen, Laura A.
 ; TITLE OF INVENTION: 22438, 23553, 25278, and 26212 No. 6534302el
 ; TITLE OF INVENTION: Human Sulfatases (A CIP Application)
 ; FILE REFERENCE: 35800/208398(5800-79
 ; CURRENT APPLICATION NUMBER: US/09/773,426A
 ; CURRENT FILING DATE: 2001-01-31
 ; PRIOR APPLICATION NUMBER: US 09/495,823
 ; PRIOR FILING DATE: 2000-01-31
 ; NUMBER OF SEQ ID NOS: 14
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO 1
 ; LENGTH: 525
 ; TYPE: PRT
 ; ORGANISM: homo sapiens
 US-09-773-426A-1

Query Match 70.5%; Score 31; DB 2; Length 525;
 Best Local Similarity 62.5%; Pred. No. 4.8e+02;
 Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IYVSLAHV 9
 :||:||:
 Db 243 LYVALAHM 250

RESULT 15

US-10-314-881-1

; Sequence 1, Application US/10314881
 ; Patent No. 6767727
 ; GENERAL INFORMATION:
 ; APPLICANT: Glucksman, Maria Alexandra
 ; APPLICANT: Williamson, Mark
 ; APPLICANT: Tsia, Fong-Ying
 ; APPLICANT: Rudolph-Owen, Laura A.
 ; TITLE OF INVENTION: 22438, 23553, 25278, and 26212 No. 6767727el
 ; TITLE OF INVENTION: Human Sulfatases (A CIP Application)
 ; FILE REFERENCE: 35800/208398(5800-79
 ; CURRENT APPLICATION NUMBER: US/10/314,881
 ; CURRENT FILING DATE: 2002-12-09

; PRIOR APPLICATION NUMBER: US/09/773,426
; PRIOR FILING DATE: 2001-01-31
; PRIOR APPLICATION NUMBER: US 09/495,823
; PRIOR FILING DATE: 2000-01-31
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 525
; TYPE: PRT
; ORGANISM: homo sapiens
US-10-314-881-1

Query Match 70.5%; Score 31; DB 2; Length 525;
Best Local Similarity 62.5%; Pred. No. 4.8e+02;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IYVSLAHV 9
:|||||:
Db 243 LYVALAHM 250

Search completed: June 30, 2008, 17:51:38
Job time : 39.625 secs

SCOPE 1.0